

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _____

Source: _____

Date Processed by STIC: _____

10/512,737A
JFL016
08/04/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

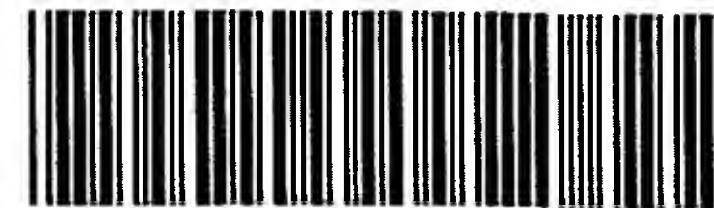
Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebs/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06



IFW16

RAW SEQUENCE LISTING

DATE: 08/04/2006

PATENT APPLICATION: US/10/512,737A

TIME: 10:13:05

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08042006\J512737A.raw

2 <110> APPLICANT: BioTeSys GmbH
 3 Schelztorstrasse 54-56
 4 D 73728 Esslingen
 5 GERMANY
 W--> 6 <120> TITLE OF INVENTION: transport system in biological systems
 C--> 7 <140> CURRENT APPLICATION NUMBER: US/10/512,737A
 C--> 7 <141> CURRENT FILING DATE: 2004-10-27
 W--> 0 <130> FILE REFERENCE:
 7 <150> PRIOR APPLICATION NUMBER: A 656/2002
 8 <151> PRIOR FILING DATE: 2002-04-29
 W--> 9 <160> NUMBER OF SEQ ID: 15

Does Not Comply
 Corrected Diskette Needed
 CP8-1-5

ERRORED SEQUENCES

W--> 10 <210> SEQ ID NO: 1
 11 <211> LENGTH: 6
 12 <212> TYPE: PRT
 13 <213> ORGANISM: Artificial sequence
 W--> 14 <220> FEATURE:
 15 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized
 W--> 16 <400> SEQUENCE: 1
 17 Gly Arg Gly Asp Ser Pro
 E--> 18 1
 19 <210> SEQ ID NO: 2
 20 <211> LENGTH: 5
 21 <212> TYPE: PRT
 22 <213> ORGANISM: Artificial sequence
 W--> 23 <220> FEATURE:
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 W--> 25 <400> SEQUENCE: 2
 26 Tyr Ile Glu Ser Arg
 E--> 27 1
 28 <210> SEQ ID NO: 3
 29 <211> LENGTH: 5
 30 <212> TYPE: PRT
 31 <213> ORGANISM: Artificial sequence
 W--> 32 <220> FEATURE:
 33 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized
 W--> 34 <400> SEQUENCE: 3
 35 Ala Asp Gly Glu Ala
 E--> 36 1
 37 <210> SEQ ID NO: 4

Invalid
 Amino Acid
 Numbering.

RAW SEQUENCE LISTING

DATE: 08/04/2006

PATENT APPLICATION: US/10/512,737A

TIME: 10:13:05

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08042006\J512737A.raw

38 <211> LENGTH: 6
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40 <213> ORGANISM: Artificial sequence
W--> 41 <220> FEATURE:
42 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized
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44 Val Arg Leu Leu Asn Asn
E--> 45 1 *S* *5*
46 <210> SEQ ID NO: 5
47 <211> LENGTH: 8
48 <212> TYPE: PRT
49 <213> ORGANISM: Artificial sequence
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51 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized
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53 Val Arg Leu Leu Asn Asn Trp Asp
E--> 54 1 *S* *5*
55 <210> SEQ ID NO: 6
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57 <212> TYPE: PRT
58 <213> ORGANISM: Artificial sequence
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60 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized
W--> 61 <400> SEQUENCE: 6
62 Gly Arg Val Arg Leu Leu Asn Asn
E--> 63 1 *S* *5*
64 <210> SEQ ID NO: 7
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66 <212> TYPE: PRT
67 <213> ORGANISM: Artificial sequence
W--> 68 <220> FEATURE:
69 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized
W--> 70 <400> SEQUENCE: 7
71 Met Thr Ala Gly Ala Gly
E--> 72 1 *S* *5*
73 <210> SEQ ID NO: 8
74 <211> LENGTH: 6
76 <212> TYPE: PRT
77 <213> ORGANISM: Artificial sequence
W--> 78 <220> FEATURE:
79 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized
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81 Leu Ser Gly Ala Leu Arg
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85 <212> TYPE: PRT
86 <213> ORGANISM: Artificial sequence
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Same error

RAW SEQUENCE LISTING

DATE: 08/04/2006

PATENT APPLICATION: US/10/512,737A

TIME: 10:13:06

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08042006\J512737A.raw

88 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized

W--> 89 <400> SEQUENCE: 9

91 Ile Val Ala Ile Leu Ile Cys Ile Leu Ile Leu Leu Thr Met Val Leu

E--> 92 1 5 5 10 10/15

E--> 93 15

94 Leu Phe Val Met Trp Met

E--> 95 20 20

96 <210> SEQ ID NO: 10

97 <211> LENGTH: 12

98 <212> TYPE: PRT

99 <213> ORGANISM: Artificial sequence

W--> 100 <220> FEATURE:

101 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized

W--> 102 <400> SEQUENCE: 10

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E--> 104 1 5 5 10 10

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106 <211> LENGTH: 18

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110 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized

W--> 111 <400> SEQUENCE: 11

112 Ile Val Ala Ile Leu Ile Cys Ile Leu Ile Leu Leu Thr Met Val Leu

E--> 113 1 5 5 10 10/15

E--> 114 15

115 Leu Phe

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117 <211> LENGTH: 6

118 <212> TYPE: PRT

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W--> 120 <220> FEATURE:

121 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized

W--> 122 <400> SEQUENCE: 12

123 Ile Val Ala Ile Leu Ile

E--> 124 1 5 5

125 <210> SEQ ID NO: 13

126 <211> LENGTH: 6

127 <212> TYPE: PRT

128 <213> ORGANISM: Artificial sequence

W--> 129 <220> FEATURE:

130 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized

W--> 131 <400> SEQUENCE: 13

132 Cys Ile Leu Ile Leu Leu

E--> 133 1 5 5

134 <210> SEQ ID NO: 14

135 <211> LENGTH: 6

136 <212> TYPE: PRT

137 <213> ORGANISM: Artificial sequence

RAW SEQUENCE LISTING

DATE: 08/04/2006

PATENT APPLICATION: US/10/512,737A

TIME: 10:13:06

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08042006\J512737A.raw

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139 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized
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141 Thr Met Val Leu Leu Phe
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143 <210> SEQ ID NO: 15
144 <211> LENGTH: 6
145 <212> TYPE: PRT
146 <213> ORGANISM: Artificial sequence
W--> 147 <220> FEATURE:
148 <223> OTHER INFORMATION: Oligopeptide Chemically Synthesized
W--> 149 <400> SEQUENCE: 15
150 Leu Phe Val Met Trp Met
E--> 151 1

Some
Error

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/512,737A

DATE: 08/04/2006
TIME: 10:13:07

Input Set : A:\pto.da.txt
Output Set: N:\CRF4\08042006\J512737A.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:9; Line(s) 92

Seq#:11; Line(s) 113

VERIFICATION SUMMARY

DATE: 08/04/2006

PATENT APPLICATION: US/10/512,737A

TIME: 10:13:07

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08042006\J512737A.raw

L:6 M:283 W: Missing Blank Line separator, <120> field identifier
 L:7 M:270 C: Current Application Number differs, Replaced Current Application No
 L:7 M:271 C: Current Filing Date differs, Replaced Current Filing Date
 L:0 M:201 W: Mandatory field data missing, <130> FILE REFERENCE
 L:9 M:283 W: Missing Blank Line separator, <160> field identifier
 L:10 M:283 W: Missing Blank Line separator, <210> field identifier
 L:14 M:283 W: Missing Blank Line separator, <220> field identifier
 L:16 M:283 W: Missing Blank Line separator, <400> field identifier
 L:18 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
 L:23 M:283 W: Missing Blank Line separator, <220> field identifier
 L:25 M:283 W: Missing Blank Line separator, <400> field identifier
 L:27 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:2
 L:32 M:283 W: Missing Blank Line separator, <220> field identifier
 L:34 M:283 W: Missing Blank Line separator, <400> field identifier
 L:36 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
 L:41 M:283 W: Missing Blank Line separator, <220> field identifier
 L:43 M:283 W: Missing Blank Line separator, <400> field identifier
 L:45 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:4
 L:50 M:283 W: Missing Blank Line separator, <220> field identifier
 L:52 M:283 W: Missing Blank Line separator, <400> field identifier
 L:54 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:5
 L:59 M:283 W: Missing Blank Line separator, <220> field identifier
 L:61 M:283 W: Missing Blank Line separator, <400> field identifier
 L:63 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
 L:68 M:283 W: Missing Blank Line separator, <220> field identifier
 L:70 M:283 W: Missing Blank Line separator, <400> field identifier
 L:72 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
 L:78 M:283 W: Missing Blank Line separator, <220> field identifier
 L:80 M:283 W: Missing Blank Line separator, <400> field identifier
 L:82 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:8
 L:87 M:283 W: Missing Blank Line separator, <220> field identifier
 L:89 M:283 W: Missing Blank Line separator, <400> field identifier
 L:92 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:9
 L:93 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:9
 L:95 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:9
 L:100 M:283 W: Missing Blank Line separator, <220> field identifier
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 L:104 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:10
 L:109 M:283 W: Missing Blank Line separator, <220> field identifier
 L:111 M:283 W: Missing Blank Line separator, <400> field identifier
 L:113 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:11
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 L:120 M:283 W: Missing Blank Line separator, <220> field identifier
 L:122 M:283 W: Missing Blank Line separator, <400> field identifier
 L:124 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:12
 L:129 M:283 W: Missing Blank Line separator, <220> field identifier
 L:131 M:283 W: Missing Blank Line separator, <400> field identifier
 L:133 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:13

VERIFICATION SUMMARY

DATE: 08/04/2006

PATENT APPLICATION: US/10/512,737A

TIME: 10:13:07

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\08042006\J512737A.raw

L:138 M:283 W: Missing Blank Line separator, <220> field identifier
L:140 M:283 W: Missing Blank Line separator, <400> field identifier
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L:147 M:283 W: Missing Blank Line separator, <220> field identifier
L:149 M:283 W: Missing Blank Line separator, <400> field identifier
L:151 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:15